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Alexander J. Blake *et al.*

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Dichloromaleic anhydride

Alexander J. Blake,* Rhiannon M. T. Griffiths, Steven M. Howdle and Claire Wilson

School of Chemistry, The University of Nottingham, University Park, Nottingham NG7 2RD, England

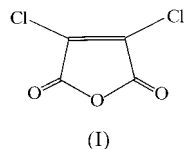
Correspondence e-mail: a.j.blake@nottingham.ac.uk

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Although molecules of the title compound, 3,4-dichloro-2,5-dihydrofuran-2,5-dione (dichloromaleic anhydride, $C_4Cl_2O_3$), (I), possess approximate non-crystallographic C_{2v} symmetry, the two chlorine substituents deviate from the ring plane. Their deviations are in the same direction, but with values of 0.0356 (17) and 0.0167 (17) Å, they differ significantly in magnitude. The closest intermolecular contact is of 2.888 (2) Å between a carbonyl O atom and the C atom of a carbonyl group, with the $O \cdots C$ direction orthogonal to the $C=O$ bond [$O5 \cdots C2^i=O2^i$ 93.6 (2)°; symmetry code: (i) $\frac{3}{2} - x, -\frac{1}{2} + y, z$]. These contacts form infinite chains of molecules running parallel to the crystallographic b direction.



Experimental

The title compound was crystallized by refluxing with 1*H*,1*H*-perfluoroheptanol.

Crystal data

$C_4Cl_2O_3$
 $M_r = 166.94$
 Orthorhombic, $Pbca$
 $a = 10.6397$ (6) Å
 $b = 8.2558$ (5) Å
 $c = 13.0057$ (7) Å
 $V = 1142.41$ (11) Å³
 $Z = 8$
 $D_x = 1.941$ Mg m⁻³

Mo $K\alpha$ radiation
 Cell parameters from 5130 reflections
 $\theta = 3.12$ – 28.55°
 $\mu = 1.050$ mm⁻¹
 $T = 150$ (2) K
 Block, colourless
 $0.32 \times 0.32 \times 0.25$ mm

Data collection

Bruker SMART CCD area-detector diffractometer
 ω scans
 Absorption correction: multi-scan (SADABS; Bruker, 1996)
 $T_{min} = 0.730$, $T_{max} = 0.779$
 7980 measured reflections
 1372 independent reflections

1261 reflections with $I > 2\sigma(I)$
 $R_{int} = 0.013$
 $\theta_{max} = 28.55^\circ$
 $h = 0 \rightarrow 14$
 $k = 0 \rightarrow 11$
 $l = 0 \rightarrow 17$
 Intensity decay: none

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.023$
 $wR(F^2) = 0.067$
 $S = 1.041$
 1372 reflections
 83 parameters

$w = 1/[\sigma^2(F_o^2) + (0.041P)^2 + 0.340P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{max} = 0.001$
 $\Delta\rho_{max} = 0.35$ e Å⁻³
 $\Delta\rho_{min} = -0.21$ e Å⁻³
 Extinction correction: SHELXL97
 Extinction coefficient: 0.0056 (7)

Data collection: SMART (Bruker, 1998); cell refinement: SMART; data reduction: SAINT (Bruker, 1999); program(s) used to solve structure: SHELXTL (Bruker, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); software used to prepare material for publication: SHELXL97 and PLATON (Spek, 1999).

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